



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/071,645

DATE: 05/30/2002

TIME: 09:06:30

Input Set : D:\Seqlist.txt

Output Set: N:\CRF3\05302002\J071645.raw

Does NOT Comply
Corrected Diskette Needed

3 <110> APPLICANT: Bihain, Bernard
 4 Bour, Barbara
 5 Bougueleret, Lydie
 7 <120> TITLE OF INVENTION: Schizophrenia Related Gene and Protein
 9 <130> FILE REFERENCE: 92.US2.CIP
 @K-> 11 <140> CURRENT APPLICATION NUMBER: US/10/071,645
 12 <141> CURRENT FILING DATE: 2002-02-06
 14 <150> PRIOR APPLICATION NUMBER: 60/223,482
 15 <151> PRIOR FILING DATE: 2000-08-07
 17 <150> PRIOR APPLICATION NUMBER: PCT/IB01/01891
 18 <151> PRIOR FILING DATE: 2001-07-26
 20 <160> NUMBER OF SEQ ID NOS: 6
 22 <170> SOFTWARE: Patent.pm

ERRORED SEQUENCES

198 <210> SEQ ID NO: 6
 199 <211> LENGTH: 592
 200 <212> TYPE: PRT
 201 <213> ORGANISM: Homo sapiens
 203 <400> SEQUENCE: 6
 204 Met Glu Thr Asp Thr Leu Leu Leu Trp Val Leu Leu Leu Trp Val Pro
 205 1 5 10 15
 206 Gly Ser Thr Gly Asp Ala Ala Gln Pro Ala Arg Arg Ala Arg Arg Thr
 207 20 25 30
 208 Tyr Glu Ala Tyr Gln Pro Leu Glu Arg Met Trp Thr Cys Asn Tyr Asn
 209 35 40 45
 210 Gln Gln Lys Asp Gln Ser Cys Asn His Lys Glu Ile Thr Ser Thr Lys
 211 50 55 60
 212 Ala Glu Arg Arg Ser Ser Gly Ile Ile Pro Val Glu Glu Glu Asn Pro
 213 65 70 75 80
 214 Asp Phe Trp Asn Arg Glu Ala Ala Glu Ala Leu Gly Ala Ala Lys Lys
 215 85 90 95
 216 Leu Gln Pro Ala Gln Thr Ala Ala Lys Asn Leu Ile Ile Phe Leu Gly
 217 100 105 110
 218 Asp Gly Met Gly Val Ser Thr Val Thr Ala Ala Arg Ile Leu Lys Gly
 219 115 120 125
 220 Gln Lys Lys Asp Lys Leu Gly Pro Glu Ile Pro Leu Ala Met Asp Arg
 221 130 135 140
 222 Phe Pro Tyr Val Ala Leu Ser Lys Thr Tyr Asn Val Asp Lys His Val
 223 145 150 155 160
 224 Pro Asp Ser Gly Ala Thr Ala Thr Ala Tyr Leu Cys Gly Val Lys Gly

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225		165		170		175
226	Asn Phe Gln Thr Ile Gly Leu Ser Ala Ala Ala Arg Phe Asn Gln Cys					
227		180		185		190
228	Asn Thr Thr Arg Gly Asn Glu Val Ile Ser Val Met Asn Arg Ala Lys					
229		195		200		205
230	Lys Ala Gly Lys Ser Val Gly Val Val Thr Thr Thr Arg Val Gln His					
231		210		215		220
232	Ala Ser Pro Ala Gly Thr Tyr Ala His Thr Val Asn Arg Asn Trp Tyr					
233	225		230		235	240
234	Ser Asp Ala Asp Val Pro Ala Ser Ala Arg Gln Glu Gly Cys Gln Asp					
235		245		250		255
236	Ile Ala Thr Gln Leu Ile Ser Asn Met Asp Ile Asp Val Ile Leu Gly					
237		260		265		270
238	Gly Gly Arg Lys Tyr Met Phe Pro Met Gly Thr Pro Asp Pro Glu Tyr					
239		275		280		285
240	Pro Asp Asp Tyr Ser Gln Gly Gly Thr Arg Leu Asp Gly Lys Asn Leu					
241		290		295		300
242	Val Gln Glu Trp Leu Ala Lys Arg Gln Gly Ala Arg Tyr Val Trp Asn					
243	305		310		315	320
244	Arg Thr Glu Leu Met Gln Ala Ser Leu Asp Pro Ser Val Thr His Leu					
245		325		330		335
246	Met Gly Leu Phe Glu Pro Gly Asp Met Lys Tyr Glu Ile His Arg Asp					
247		340		345		350
248	Ser Thr Leu Asp Pro Ser Leu Met Glu Met Thr Glu Ala Ala Leu Arg					
249		355		360		365
250	Leu Leu Ser Arg Asn Pro Arg Gly Phe Phe Leu Phe Val Glu Gly Gly					
251		370		375		380
252	Arg Ile Asp His Gly His His Glu Ser Arg Ala Tyr Arg Ala Leu Thr					
253	385		390		395	400
254	Glu Thr Ile Met Phe Asp Asp Ala Ile Glu Arg Ala Gly Gln Leu Thr					
255		405		410		415
256	Ser Glu Glu Asp Thr Leu Ser Leu Val Thr Ala Asp His Ser His Val					
257		420		425		430
258	Phe Ser Phe Gly Gly Tyr Pro Leu Arg Gly Ser Ser Ile Phe Gly Leu					
259		435		440		445
260	Ala Pro Gly Lys Ala Arg Asp Arg Lys Ala Tyr Thr Val Leu Leu Tyr					
261		450		455		460
262	Gly Asn Gly Pro Gly Tyr Val Leu Lys Asp Gly Ala Arg Pro Asp Val					
263	465		470		475	480
264	Thr Glu Ser Glu Ser Gly Ser Pro Glu Tyr Arg Gln Gln Ser Ala Val					
265		485		490		495
266	Pro Leu Asp Glu Glu Thr His Ala Gly Glu Asp Val Ala Val Phe Ala					
267		500		505		510
268	Arg Gly Pro Gln Ala His Leu Val His Gly Val Gln Glu Gln Thr Phe					
269		515		520		525
270	Ile Ala His Val Met Ala Phe Ala Ala Cys Leu Glu Pro Tyr Thr Ala					
271		530		535		540
272	Cys Asp Leu Ala Pro Pro Ala Gly Thr Thr Asp Ala Ala His Pro Gly					
273	545		550		555	560

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274 Tyr Leu Glu Glu Ala Leu Ser Leu Glu Gly Pro Glu Gln Lys Leu Ile
275 565 570 575
276 Ser Glu Glu Asp Leu Asn Ser Ala Val Asp His His His His His His
277 580 585 590
E--> 282 (1)

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/071,645

DATE: 05/30/2002

TIME: 09:06:32

Input Set : D:\Seqlist.txt

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L:11 M:270 C: Current Application Number differs, Replaced Application Number

L:282 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6